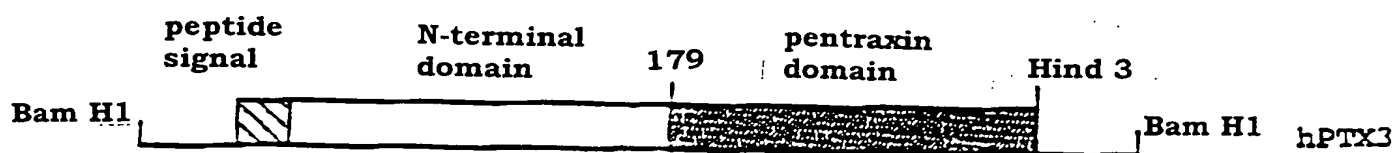


FIGURE 1



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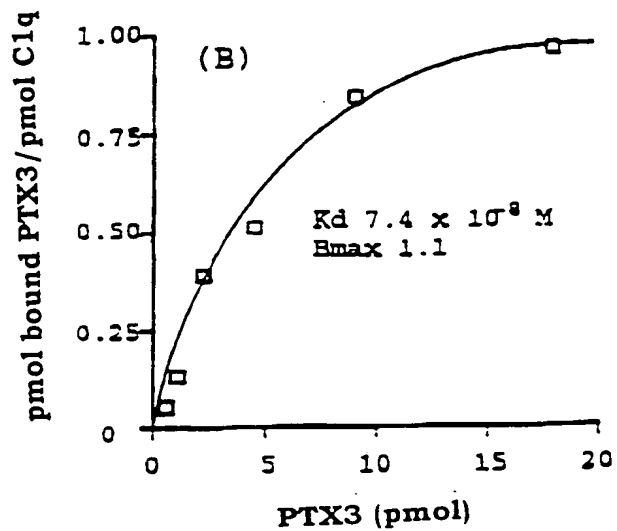
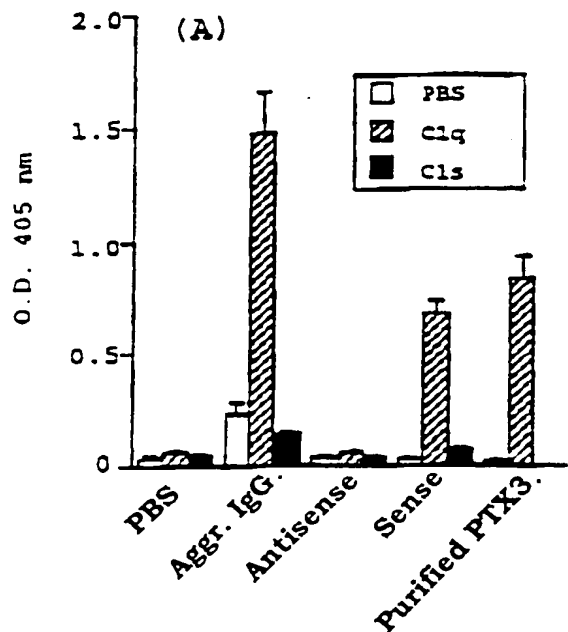
FIGURE 2

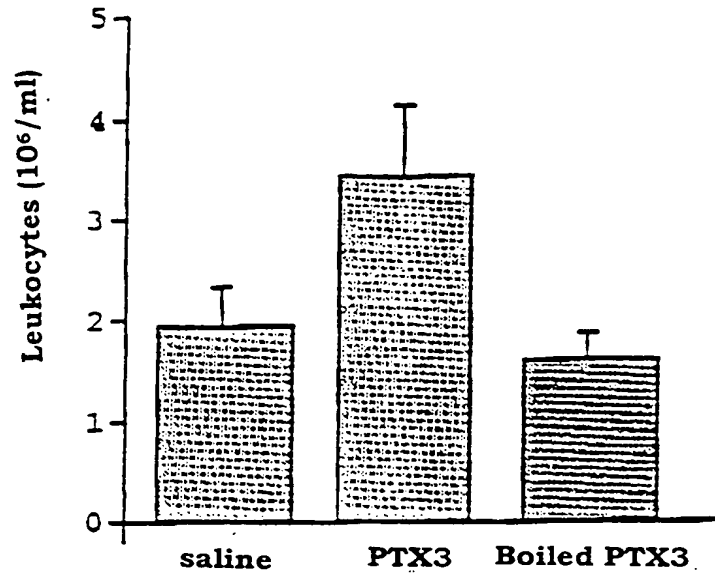
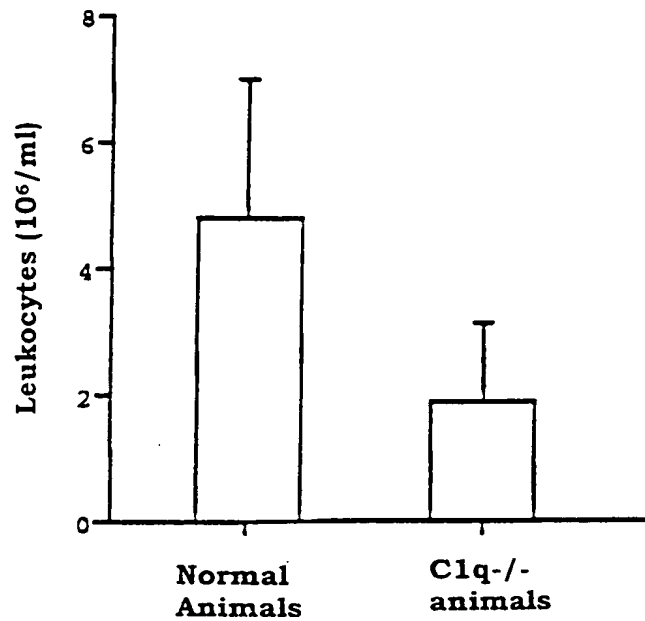
FIGURE 3

FIGURE 4

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SEQ. ID. NO:

SequenceV1

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RGELQRLREE	LGRLAESLAR	PCAPGAPAEA	RLTSALDELL			121
QATSDAGRRL	ARMEGAFAQR	PEEAGRALAA	VLEELRQTRA			161
DLFAVQGWAA	RSWLFAGCET	AILFPMRSKK	IFGSVHPVRP			201
MRLESFSACI	WVKATDVLNK	TILFSYGTKR	NPYEIQLYLS			241
YQSIIVFVGG	EENKLVAEAM	VSLGRWTHLC	GTWNSEEGLT			281
SLWVNGELAA	TTVEMATGHI	VPEGGILQIG	QEKNGCCVGG			321
GFDETLAFSG	RLTGFNWDS	VLSNZEIRET	GGAESCHIRG			361
NIVGWGVTEI	QPHGGAQYVS					381

004550 004550 004550

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SEQ. ID. NO:

Sequence 2

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